SEQUENCE LISTING

<110> Ono Pharmaceutical Co., Ltd.

<120> A novel polypeptide, a cDNA encoding the polypeptide and utilization thereof

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<151> 1999-04-28

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Val Leu Leu Asp Asp Asn Arg Ser Cys Gln Asp Ile Asn Glu Cys Glu
255                 260                 265                 270

cac cga aac cac acg tgt acc tca ctg cag act tgc tac aat cta caa   1197
His Arg Asn His Thr Cys Thr Ser Leu Gln Thr Cys Tyr Asn Leu Gln
                275                 280                 285
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ggg ggc ttc aaa tgt att gat ccc atc agc tgt gag gag cct tat ctg   1245
Gly Gly Phe Lys Cys Ile Asp Pro Ile Ser Cys Glu Glu Pro Tyr Leu
            290                 295                 300

ctg att ggt gaa aac cgc tgt atg tgt cct gct gag cac acc agc tgc   1293
Leu Ile Gly Glu Asn Arg Cys Met Cys Pro Ala Glu His Thr Ser Cys
        305                 310                 315

aga gac cag cca ttc acc atc ctg tat cgg gac atg gat gtg gtg tca   1341
Arg Asp Gln Pro Phe Thr Ile Leu Tyr Arg Asp Met Asp Val Val Ser
    320                 325                 330

gga cgc tcc gtt cct gct gac atc ttc cag atg caa gca aca acc cga   1389
Gly Arg Ser Val Pro Ala Asp Ile Phe Gln Met Gln Ala Thr Thr Arg
335                 340                 345                 350

tac cct ggt gcc tat tac att ttc cag atc aaa tct ggc aac gag ggt   1437
Tyr Pro Gly Ala Tyr Tyr Ile Phe Gln Ile Lys Ser Gly Asn Glu Gly
                355                 360                 365

cga gag ttc tat atg cgg caa aca ggg cct atc agt gcc acc ctg gtg   1485
Arg Glu Phe Tyr Met Arg Gln Thr Gly Pro Ile Ser Ala Thr Leu Val
            370                 375                 380

atg aca cgc ccc atc aaa ggg cct cgg gac atc cag ctg gac ttg gag   1533
Met Thr Arg Pro Ile Lys Gly Pro Arg Asp Ile Gln Leu Asp Leu Glu
        385                 390                 395

atg atc act gtc aac act gtc atc aac ttc aga ggc agc tcc gtg atc   1581
Met Ile Thr Val Asn Thr Val Ile Asn Phe Arg Gly Ser Ser Val Ile
    400                 405                 410

cga ctg cgg ata tat gtg tcg cag tat ccg ttc tgagcctctg gctaaggcct 1634
Arg Leu Arg Ile Tyr Val Ser Gln Tyr Pro Phe
415                 420                 425

ctgacactgc ctttcaccag caccgaggga cgggaggaga aaggaaacca gcaagaatga 1694

gagcgagaca gacattgcac ctttcctgct gaatatctcc tgggggcatc agcctagcat 1754

cttgacccat atctgtacta ttgcagatgg tcactctgaa ggacaccctg ccctcagttc 1814

ctatgatgca gttatccaaa agtgttcatc ttagccccctg atatgaggtt gccagtgact 1874

cttcaaagcc ttccatttat ttccatcgtt ttataaaaaa gaaaatagat tagatttgct 1934

ggggtatgag tcctcgaagg ttcaaaagac tgagtggctt gctctcacct cttcctctcc 1994
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ttcctccatc tcttgctgca ttgctgcttt gcaaaagtcc tcatgggctc gtgggaaatg 2054

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aggatcgaag gtttttatag agtctatttt aaaatcacat ctggtatttt cagcataaaa 2174

gaaattttag ttgtctttaa aatttgtatg agtgtttaac cttttcttat tcattttgag 2234

gcttcttaaa gtggtagaat tccttccaaa ggcctcagat acatgttatg ttcagtcttt 2294

ccaacctcat cctttcctgc atcttagccc agtttttacg aagaccccct aatcatgctt 2354

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tgaagaaaaa aaaaa 2429

<210> 8
<211> 461
<212> PRT
<213> Mus musculus
<223> Clone mouse A55b derived from Day 13 mouse
embryonic heart

<400> 8

Met	Gly	Pro	Arg	Ser	Phe	Glu	Pro	Met	His	Ser	Gly	Leu	Cys	Arg	Gln
-35						-30					-25				
Arg	Arg	Met	Ile	Leu	Thr	Val	Thr	Ile	Leu	Ala	Leu	Trp	Leu	Pro	His
-20					-15					-10					-5
Pro	Gly	Asn	Ala	Gln	Gln	Gln	Cys	Thr	Asn	Gly	Phe	Asp	Leu	Asp	Arg
			-1	1				5					10		
Gln	Ser	Gly	Gln	Cys	Leu	Asp	Ile	Asp	Glu	Cys	Arg	Thr	Ile	Pro	Glu
		15					20					25			
Ala	Cys	Arg	Gly	Asp	Met	Met	Cys	Val	Asn	Gln	Asn	Gly	Gly	Tyr	Leu
	30					35					40				
Cys	Ile	Pro	Arg	Thr	Asn	Pro	Val	Tyr	Arg	Gly	Pro	Tyr	Ser	Asn	Pro
45					50					55					60
Tyr	Ser	Thr	Ser	Tyr	Ser	Gly	Pro	Tyr	Pro	Ala	Ala	Ala	Pro	Pro	Val
			65					70					75		
Pro	Ala	Ser	Asn	Tyr	Pro	Thr	Ile	Ser	Arg	Pro	Leu	Val	Cys	Arg	Phe
		80					85					90			

Gly Tyr Gln Met Asp Glu Gly Asn Gln Cys Val Asp Val Asp Glu Cys
95 100 105

Ala Thr Asp Ser His Gln Cys Asn Pro Thr Gln Ile Cys Ile Asn Thr
110 115 120

Glu Gly Gly Tyr Thr Cys Ser Cys Thr Asp Gly Tyr Trp Leu Leu Glu
125 130 135 140

Gly Gln Cys Leu Asp Ile Asp Glu Cys Arg Tyr Gly Tyr Cys Gln Gln
145 150 155

Leu Cys Ala Asn Val Pro Gly Ser Tyr Ser Cys Thr Cys Asn Pro Gly
160 165 170

Phe Thr Leu Asn Asp Asp Gly Arg Ser Cys Gln Asp Val Asn Glu Cys
175 180 185

Glu Thr Glu Asn Pro Cys Val Gln Thr Cys Val Asn Thr Tyr Gly Ser
190 195 200

Phe Ile Cys Arg Cys Asp Pro Gly Tyr Glu Leu Glu Glu Asp Gly Ile
205 210 215 220

His Cys Ser Asp Met Asp Glu Cys Ser Phe Ser Glu Phe Leu Cys Gln
225 230 235

His Glu Cys Val Asn Gln Pro Gly Ser Tyr Phe Cys Ser Cys Pro Pro
240 245 250

Gly Tyr Val Leu Leu Asp Asp Asn Arg Ser Cys Gln Asp Ile Asn Glu
255 260 265

Cys Glu His Arg Asn His Thr Cys Thr Ser Leu Gln Thr Cys Tyr Asn
270 275 280

Leu Gln Gly Gly Phe Lys Cys Ile Asp Pro Ile Ser Cys Glu Glu Pro
285 290 295 300

Tyr Leu Leu Ile Gly Glu Asn Arg Cys Met Cys Pro Ala Glu His Thr
305 310 315

Ser Cys Arg Asp Gln Pro Phe Thr Ile Leu Tyr Arg Asp Met Asp Val
320 325 330

Val Ser Gly Arg Ser Val Pro Ala Asp Ile Phe Gln Met Gln Ala Thr
335 340 345

Thr Arg Tyr Pro Gly Ala Tyr Tyr Ile Phe Gln Ile Lys Ser Gly Asn
350 355 360

Glu Gly Arg Glu Phe Tyr Met Arg Gln Thr Gly Pro Ile Ser Ala Thr
365 370 375 380

Leu Val Met Thr Arg Pro Ile Lys Gly Pro Arg Asp Ile Gln Leu Asp
385 390 395

Leu Glu Met Ile Thr Val Asn Thr Val Ile Asn Phe Arg Gly Ser Ser
400 405 410

Val Ile Arg Leu Arg Ile Tyr Val Ser Gln Tyr Pro Phe
415 420 425

<210> 9
<211> 423
<212> PRT
<213> Mus musculus

<400> 9
Gln Cys Thr Asn Gly Phe Asp Leu Asp Arg Gln Ser Gly Gln Cys Leu
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Asp Ile Asp Glu Cys Arg Thr Ile Pro Glu Ala Cys Arg Gly Asp Met
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Met Cys Val Asn Gln Asn Gly Gly Tyr Leu Cys Ile Pro Arg Thr Asn
35 40 45

Pro Val Tyr Arg Gly Pro Tyr Ser Asn Pro Tyr Ser Thr Ser Tyr Ser
50 55 60

Gly Pro Tyr Pro Ala Ala Ala Pro Pro Val Pro Ala Ser Asn Tyr Pro
65 70 75 80

Thr Ile Ser Arg Pro Leu Val Cys Arg Phe Gly Tyr Gln Met Asp Glu
85 90 95

Gly Asn Gln Cys Val Asp Val Asp Glu Cys Ala Thr Asp Ser His Gln
100 105 110

Cys Asn Pro Thr Gln Ile Cys Ile Asn Thr Glu Gly Gly Tyr Thr Cys
115 120 125

Ser Cys Thr Asp Gly Tyr Trp Leu Leu Glu Gly Gln Cys Leu Asp Ile
130 135 140

Asp Glu Cys Arg Tyr Gly Tyr Cys Gln Gln Leu Cys Ala Asn Val Pro
145 150 155 160

Gly Ser Tyr Ser Cys Thr Cys Asn Pro Gly Phe Thr Leu Asn Asp Asp
165 170 175

Gly Arg Ser Cys Gln Asp Val Asn Glu Cys Glu Thr Glu Asn Pro Cys
180 185 190

Val Gln Thr Cys Val Asn Thr Tyr Gly Ser Phe Ile Cys Arg Cys Asp
195 200 205

Pro Gly Tyr Glu Leu Glu Glu Asp Gly Ile His Cys Ser Asp Met Asp
210 215 220

Glu Cys Ser Phe Ser Glu Phe Leu Cys Gln His Glu Cys Val Asn Gln
225 230 235 240

Pro Gly Ser Tyr Phe Cys Ser Cys Pro Pro Gly Tyr Val Leu Leu Asp
245 250 255

Asp Asn Arg Ser Cys Gln Asp Ile Asn Glu Cys Glu His Arg Asn His
260 265 270

Thr Cys Thr Ser Leu Gln Thr Cys Tyr Asn Leu Gln Gly Gly Phe Lys
275 280 285

Cys Ile Asp Pro Ile Ser Cys Glu Glu Pro Tyr Leu Leu Ile Gly Glu
290 295 300

Asn Arg Cys Met Cys Pro Ala Glu His Thr Ser Cys Arg Asp Gln Pro
305 310 315 320

Phe Thr Ile Leu Tyr Arg Asp Met Asp Val Val Ser Gly Arg Ser Val
325 330 335

Pro Ala Asp Ile Phe Gln Met Gln Ala Thr Thr Arg Tyr Pro Gly Ala
340 345 350

Tyr Tyr Ile Phe Gln Ile Lys Ser Gly Asn Glu Gly Arg Glu Phe Tyr
355 360 365

Met Arg Gln Thr Gly Pro Ile Ser Ala Thr Leu Val Met Thr Arg Pro
370 375 380

Ile Lys Gly Pro Arg Asp Ile Gln Leu Asp Leu Glu Met Ile Thr Val
385 390 395 400

Asn Thr Val Ile Asn Phe Arg Gly Ser Ser Val Ile Arg Leu Arg Ile
405 410 415

Tyr Val Ser Gln Tyr Pro Phe
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<210> 10
<211> 1269
<212> DNA
<213> Mus musculus

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tatttgtgca tccctcgaac caacccagtg tatcgagggc cttactcaaa tccctactct 180
acatcctact caggcccata cccagcagcg gccccaccag taccagcttc caactacccc 240
acgatttcaa ggcctcttgt ctgccgcttt gggtatcaga tggatgaagg caaccagtgt 300
gtggatgtgg acgagtgtgc aacagactca caccagtgca accctaccca gatctgtatc 360
aacactgaag gaggttacac ctgctcctgc accgatgggt actggcttct ggaagggcag 420
tgcctagata ttgatgaatg tcgctatggt tactgccagc agctctgtgc aaatgttcca 480
ggatcctatt cctgtacatg caaccctggt ttcaccctca acgacgatgg aaggtcttgc 540
caagatgtga acgagtgcga aactgagaat ccctgtgttc agacctgtgt caacacctat 600
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agtgatatgg acgagtgcag cttctccgag ttcctctgtc aacacgagtg tgtgaaccag 720
ccgggctcat acttctgctc gtgccctcca ggctacgtcc tgttggatga taaccgaagc 780
tgccaggata tcaatgaatg tgagcaccga aaccacacgt gtacctcact gcagacttgc 840
tacaatctac aagggggctt caaatgtatt gatcccatca gctgtgagga gccttatctg 900
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ttccagatgc aagcaacaac ccgataccct ggtgcctatt acatttttcca gatcaaatct 1080
ggcaacgagg gtcgagagtt ctatatgcgg caaacagggc ctatcagtgc caccctggtg 1140
atgacacgcc ccatcaaagg gcctcgggac atccagctgg acttggagat gatcactgtc 1200
aacactgtca tcaacttcag aggcagctcc gtgatccgac tgcggatata tgtgtcgcag 1260
tatccgttc 1269

<210> 11
<211> 35
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 11
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<210> 12
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: mA55 R1 Primer

<400> 12
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